

qPCR analysis of selected insertions and spike-in controls

Locus	Unnorm*	Norm1*	Norm2*
FI-11a	10.5 (0.14)	14.0 (0.22)	15.5 (0.02)
FI-14a	11.7 (0.21)	13.9 (0.22)	14.8 (0.12)
FI-18a	11.0 (0.06)	15.8 (0.19)	15.6 (0.06)
DR240	28.6 (0.14)	23.7 (0.29)	21.1 (0.07)
DR389	26.2 (0.10)	23.3 (0.32)	21.4 (0.04)
DR418	32.9 (0.73)	25.8 (0.14)	24.9 (0.09)
DR259	35.2 (0.2)	27.0 (0.44)	23.8 (0.02)
SI-1	22.2 (0.11)	19.9 (0.06)	21.4 (0.06)
SI-3	21.3 (0.04)	18.9 (0.08)	17.5 (0.05)
SI-13	22.4 (0.18)	20.0 (0.07)	18.8 (0.13)
SI-17	24.1 (0.19)	21.5 (0.13)	20.7 (0.12)

\* - for each insertion Ct and standard deviation (in parentheses) is given

FI – fixed insertion present in each cell analyzed, DR – spike-in controls, SI – somatic insertions.